Serial Number: 09/847,208

CRF Processing Date: 5/16/2001
Edited by: _A_
Verified by: _A_ _____ (STIC staff)

ENTERED

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: .
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING   DATE: 05/16/2001
PATENT APPLICATION: US/09/847,208   TIME: 15:34:47

Input Set : **A:\DES.txt**
Output Set: **N:\CRF3\05162001\I847208.raw**

```
     3 <110> APPLICANT: Saxon, Andrew
     4        Zhang, Ke
     5        Zhu, Daocheng
     7 <120> TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
     8        IgE-MEDIATED ALLERGIC DISEASES
    10 <130> FILE REFERENCE: UC67.002A
C-->12 <140> CURRENT APPLICATION NUMBER: US/09/847,208
C-->12 <141> CURRENT FILING DATE: 2001-05-01
    12 <160> NUMBER OF SEQ ID NOS: 177
    14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
```

ERRORED SEQUENCES

```
  6232 <210> SEQ ID NO: 177
  6233 <211> LENGTH: 32
  6234 <212> TYPE: PRT
  6235 <213> ORGANISM: Homo sapiens
  6237 <220> FEATURE:
  6238 <221> NAME/KEY: UNSURE
  6239 <222> LOCATION: 1, 4, 8, 10, 11, 12, 27, 30
  6240 <223> OTHER INFORMATION: Xaa = any amino acid
  6242 <400> SEQUENCE: 177
W-->6243 Xaa Gln Gln Xaa Glu Leu Gln Xaa Asp Xaa Xaa Xaa Gln Ser Gln Leu
  6244  1               5                 10                  15
W-->6245 Glu Arg Ala Asp Leu Arg Pro Gly Glu Gln Xaa Leu Met Xaa Lys Ile
  6246            20                25                  30
E-->6251 W:\DOCS\BGY\SEQUENCE LISTING\UC67.003A.doc
E-->6252 5
E-->6254 1
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/847,208

DATE: 05/16/2001
TIME: 15:34:49

Input Set : **A:\DES.txt**
Output Set: **N:\CRF3\05162001\I847208.raw**

```
L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:3086 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:3129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:3466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93
L:3468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93
L:3483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94
L:4518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:126
L:6226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176
L:6228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176
L:6243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177
L:6245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177
L:6251 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:6251 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:6252 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:177
M:332 Repeated in SeqNo=177
L:6254 M:252 E: No. of Seq. differs, <211>LENGTH:Input:32 Found:34 SEQ:177
```